

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/526,978
Source: P4710
Date Processed by STIC: 3/16/05

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 03/16/2005

PATENT APPLICATION: US/10/526,978

TIME: 08:11:26

Input Set : A:\Ms0023p.txt

Output Set: N:\CRF4\03162005\J526978.raw

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4 <110> APPLICANT: Daggett, Lorrie
5      Morales, Sylvia
6      Schaffhauser, Herve
8 <120> TITLE OF INVENTION: NUCLEIC ACID SEQUENCES ENCODING NOVEL
9      POINT MUTATIONS ON mGluR2 AND mGluR3, POLYPEPTIDES WITH SAID
10     MUTATIONS, AND METHODS OF USING SAID NUCLEIC ACID SEQUENCES,
11     AND SAID POLYPEPTIDES, TO IDENTIFY, PREDICT AND EVALUATE
12     SPECIFIC, SELECTIVE MODULATORS WHOSE ASSOCIATION TO mGluR2
13     OR mGluR3 IS EFFECTED BY SAID MUTATIONS
16 <130> FILE REFERENCE: MS0023P
C--> 18 <140> CURRENT APPLICATION NUMBER: US/10/526,978
C--> 18 <141> CURRENT FILING DATE: 2005-03-07
18 <150> PRIOR APPLICATION NUMBER: PCT/US03/28470
19 <151> PRIOR FILING DATE: 2003-09-11
21 <150> PRIOR APPLICATION NUMBER: 60/409,910
22 <151> PRIOR FILING DATE: 2002-09-11
24 <160> NUMBER OF SEQ ID NOS: 48
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 873
30 <212> TYPE: PRT
31 <213> ORGANISM: Human
33 <400> SEQUENCE: 1
34 Met Gly Ser Leu Leu Ala Leu Leu Ala Leu Leu Pro Leu Trp Gly Ala
35 1          5          10          15
36 Val Ala Glu Gly Pro Ala Lys Lys Val Leu Thr Leu Glu Gly Asp Leu
37 20          25          30
38 Val Leu Gly Gly Leu Phe Pro Val His Gln Lys Gly Gly Pro Ala Glu
39 35          40          45
40 Asp Cys Gly Pro Val Asn Glu His Arg Gly Ile Gln Arg Leu Glu Ala
41 50          55          60
42 Met Leu Phe Ala Leu Asp Arg Ile Asn Arg Asp Pro His Leu Leu Pro
43 65          70          75          80
44 Gly Val Arg Leu Gly Ala His Ile Leu Asp Ser Cys Ser Lys Asp Thr
45 85          90          95
46 His Ala Leu Glu Gln Ala Leu Asp Phe Val Arg Ala Ser Leu Ser Arg
47 100         105         110
48 Gly Ala Asp Gly Ser Arg His Ile Cys Pro Asp Gly Ser Tyr Ala Thr
49 115         120         125
50 His Gly Asp Ala Pro Thr Ala Ile Thr Gly Val Ile Gly Gly Ser Tyr
51 130         135         140
52 Ser Asp Val Ser Ile Gln Val Ala Asn Leu Leu Arg Leu Phe Gln Ile
53 145         150         155         160

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54 Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala Lys Leu Ser Asp Lys Ser
55           165           170           175
56 Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro Pro Asp Phe Phe Gln Ala
57           180           185           190
58 Lys Ala Met Ala Glu Ile Leu Arg Phe Phe Asn Trp Thr Tyr Val Ser
59           195           200           205
60 Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu Thr Gly Ile Glu Ala Phe
61           210           215           220
62 Glu Leu Glu Ala Arg Ala Arg Asn Ile Cys Val Ala Thr Ser Glu Lys
63 225           230           235           240
64 Val Gly Arg Ala Met Ser Arg Ala Ala Phe Glu Gly Val Val Arg Ala
65           245           250           255
66 Leu Leu Gln Lys Pro Ser Ala Arg Val Ala Val Leu Phe Thr Arg Ser
67           260           265           270
68 Glu Asp Ala Arg Glu Leu Leu Ala Ala Ser Gln Arg Leu Asn Ala Ser
69           275           280           285
70 Phe Thr Trp Val Ala Ser Asp Gly Trp Gly Ala Leu Glu Ser Val Val
71           290           295           300
72 Ala Gly Ser Glu Gly Ala Ala Glu Gly Ala Ile Thr Ile Glu Leu Ala
73 305           310           315           320
74 Ser Tyr Pro Ile Ser Asp Phe Ala Ser Tyr Phe Gln Ser Leu Asp Pro
75           325           330           335
76 Trp Asn Asn Ser Arg Asn Pro Trp Phe Arg Glu Phe Trp Glu Gln Arg
77           340           345           350
78 Phe Arg Cys Ser Phe Arg Gln Arg Asp Cys Ala Ala His Ser Leu Arg
79           355           360           365
80 Ala Val Pro Phe Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala
81           370           375           380
82 Val Tyr Ala Met Ala His Ala Leu His Asn Met His Arg Ala Leu Cys
83 385           390           395           400
84 Pro Asn Thr Thr Arg Leu Cys Asp Ala Met Arg Pro Val Asn Gly Arg
85           405           410           415
86 Arg Leu Tyr Lys Asp Phe Val Leu Asn Val Lys Phe Asp Ala Pro Phe
87           420           425           430
88 Arg Pro Ala Asp Thr His Asn Glu Val Arg Phe Asp Arg Phe Gly Asp
89           435           440           445
90 Gly Ile Gly Arg Tyr Asn Ile Phe Thr Tyr Leu Arg Ala Gly Ser Gly
91           450           455           460
92 Arg Tyr Arg Tyr Gln Lys Val Gly Tyr Trp Ala Glu Gly Leu Thr Leu
93 465           470           475           480
94 Asp Thr Ser Leu Ile Pro Trp Ala Ser Pro Ser Ala Gly Pro Leu Pro
95           485           490           495
96 Ala Ser Arg Cys Ser Glu Pro Cys Leu Gln Asn Glu Val Lys Ser Val
97           500           505           510
98 Gln Pro Gly Glu Val Cys Cys Trp Leu Cys Ile Pro Cys Gln Pro Tyr
99           515           520           525
100 Glu Tyr Arg Leu Asp Glu Phe Thr Cys Ala Asp Cys Gly Leu Gly Tyr
101           530           535           540
102 Trp Pro Asn Ala Ser Leu Thr Gly Cys Phe Glu Leu Pro Gln Glu Tyr

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103 545          550          555          560
104 Ile Arg Trp Gly Asp Ala Trp Ala Val Gly Pro Val Thr Ile Ala Cys
105          565          570          575
106 Leu Gly Ala Leu Ala Thr Leu Phe Val Leu Gly Val Phe Val Arg His
107          580          585          590
108 Asn Ala Thr Pro Val Val Lys Ala Ser Gly Arg Glu Leu Cys Tyr Ile
109          595          600          605
110 Leu Leu Gly Gly Val Phe Leu Cys Tyr Cys Met Thr Phe Ile Phe Ile
111          610          615          620
112 Ala Lys Pro Ser Thr Ala Val Cys Thr Leu Arg Arg Leu Gly Leu Gly
113 625          630          635          640
114 Thr Ala Phe Ser Val Cys Tyr Ser Ala Leu Leu Thr Lys Thr Asn Arg
115          645          650          655
116 Ile Ala Arg Ile Phe Gly Gly Ala Arg Glu Gly Ala Gln Arg Pro Arg
117          660          665          670
118 Phe Ile Ser Pro Ala Ser Gln Val Ala Ile Cys Leu Ala Leu Ile Ser
119          675          680          685
120 Gly Gln Leu Leu Ile Val Val Ala Trp Leu Val Val Glu Ala Pro Gly
121          690          695          700
122 Thr Gly Lys Glu Thr Ala Pro Glu Arg Arg Glu Val Val Thr Leu Arg
123 705          710          715          720
124 Cys Asn His Arg Asp Ala Ser Met Leu Gly Ser Leu Ala Tyr Asp Val
125          725          730          735
126 Leu Leu Ile Ala Leu Cys Thr Leu Tyr Ala Phe Lys Thr Arg Lys Cys
127          740          745          750
128 Pro Glu Asn Phe Asn Glu Ala Lys Phe Ile Gly Phe Thr Met Tyr Thr
129          755          760          765
130 Thr Cys Ile Ile Trp Leu Ala Phe Leu Pro Ile Phe Tyr Val Thr Ser
131          770          775          780
132 Ser Asp Tyr Arg Val Gln Thr Thr Thr Met Cys Val Ser Val Ser Leu
133 785          790          795          800
134 Ser Gly Ser Val Val Leu Gly Cys Leu Phe Ala Pro Lys Leu His Ile
135          805          810          815
136 Ile Leu Phe Gln Pro Gln Lys Asn Val Val Ser His Arg Ala Pro Thr
137          820          825          830
138 Ser Arg Phe Gly Ser Ala Ala Ala Arg Ala Ser Ser Ser Leu Gly Gln
139          835          840          845
140 Gly Ser Gly Ser Gln Phe Val Pro Thr Val Cys Asn Gly Arg Glu Val
141          850          855          860
142 Val Asp Ser Thr Thr Ser Ser Leu Thr
143 865          870
146 <210> SEQ ID NO: 2
147 <211> LENGTH: 873
148 <212> TYPE: PRT
149 <213> ORGANISM: Human
151 <400> SEQUENCE: 2
152 Met Gly Ser Leu Leu Ala Leu Leu Ala Leu Leu Pro Leu Trp Gly Ala
153 1          5          10          15
154 Val Ala Glu Gly Pro Ala Lys Lys Val Leu Thr Leu Glu Gly Asp Leu

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155			20				25			30						
156	Val	Leu	Gly	Gly	Leu	Phe	Pro	Val	His	Gln	Lys	Gly	Gly	Pro	Ala	Glu
157			35				40					45				
158	Asp	Cys	Gly	Pro	Val	Asn	Glu	His	Arg	Gly	Ile	Gln	Arg	Leu	Glu	Ala
159		50					55					60				
160	Met	Leu	Phe	Ala	Leu	Asp	Arg	Ile	Asn	Arg	Asp	Pro	His	Leu	Leu	Pro
161	65					70					75					80
162	Gly	Val	Arg	Leu	Gly	Ala	His	Ile	Leu	Asp	Ser	Cys	Ser	Lys	Asp	Thr
163				85						90					95	
164	His	Ala	Leu	Glu	Gln	Ala	Leu	Asp	Phe	Val	Arg	Ala	Ser	Leu	Ser	Arg
165				100					105						110	
166	Gly	Ala	Asp	Gly	Ser	Arg	His	Ile	Cys	Pro	Asp	Gly	Ser	Tyr	Ala	Thr
167			115					120						125		
168	His	Gly	Asp	Ala	Pro	Thr	Ala	Ile	Thr	Gly	Val	Ile	Gly	Gly	Ser	Tyr
169		130					135					140				
170	Ser	Asp	Val	Ser	Ile	Gln	Val	Ala	Asn	Leu	Leu	Arg	Leu	Phe	Gln	Ile
171	145					150					155					160
172	Pro	Gln	Ile	Ser	Tyr	Ala	Ser	Thr	Ser	Ala	Lys	Leu	Ser	Asp	Lys	Ser
173					165					170					175	
174	Arg	Tyr	Asp	Tyr	Phe	Ala	Arg	Thr	Val	Pro	Pro	Asp	Phe	Phe	Gln	Ala
175				180					185						190	
176	Lys	Ala	Met	Ala	Glu	Ile	Leu	Arg	Phe	Phe	Asn	Trp	Thr	Tyr	Val	Ser
177			195					200						205		
178	Thr	Val	Ala	Ser	Glu	Gly	Asp	Tyr	Gly	Glu	Thr	Gly	Ile	Glu	Ala	Phe
179		210					215					220				
180	Glu	Leu	Glu	Ala	Arg	Ala	Arg	Asn	Ile	Cys	Val	Ala	Thr	Ser	Glu	Lys
181	225					230					235					240
182	Val	Gly	Arg	Ala	Met	Ser	Arg	Ala	Ala	Phe	Glu	Gly	Val	Val	Arg	Ala
183				245						250					255	
184	Leu	Leu	Gln	Lys	Pro	Ser	Ala	Arg	Val	Ala	Val	Leu	Phe	Thr	Arg	Ser
185				260					265						270	
186	Glu	Asp	Ala	Arg	Glu	Leu	Leu	Ala	Ala	Ser	Gln	Arg	Leu	Asn	Ala	Ser
187			275					280					285			
188	Phe	Thr	Trp	Val	Ala	Ser	Asp	Gly	Trp	Gly	Ala	Leu	Glu	Ser	Val	Val
189		290					295					300				
190	Ala	Gly	Ser	Glu	Gly	Ala	Ala	Glu	Gly	Ala	Ile	Thr	Ile	Glu	Leu	Ala
191	305					310					315					320
192	Ser	Tyr	Pro	Ile	Ser	Asp	Phe	Ala	Ser	Tyr	Phe	Gln	Ser	Leu	Asp	Pro
193					325					330					335	
194	Trp	Asn	Asn	Ser	Arg	Asn	Pro	Trp	Phe	Arg	Glu	Phe	Trp	Glu	Gln	Arg
195				340					345						350	
196	Phe	Arg	Cys	Ser	Phe	Arg	Gln	Arg	Asp	Cys	Ala	Ala	His	Ser	Leu	Arg
197			355					360					365			
198	Ala	Val	Pro	Phe	Glu	Gln	Glu	Ser	Lys	Ile	Met	Phe	Val	Val	Asn	Ala
199		370					375					380				
200	Val	Tyr	Ala	Met	Ala	His	Ala	Leu	His	Asn	Met	His	Arg	Ala	Leu	Cys
201	385					390					395					400
202	Pro	Asn	Thr	Thr	Arg	Leu	Cys	Asp	Ala	Met	Arg	Pro	Val	Asn	Gly	Arg
203					405					410					415	

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204 Arg Leu Tyr Lys Asp Phe Val Leu Asn Val Lys Phe Asp Ala Pro Phe
205          420          425          430
206 Arg Pro Ala Asp Thr His Asn Glu Val Arg Phe Asp Arg Phe Gly Asp
207          435          440          445
208 Gly Ile Gly Arg Tyr Asn Ile Phe Thr Tyr Leu Arg Ala Gly Ser Gly
209          450          455          460
210 Arg Tyr Arg Tyr Gln Lys Val Gly Tyr Trp Ala Glu Gly Leu Thr Leu
211 465          470          475          480
212 Asp Thr Ser Leu Ile Pro Trp Ala Ser Pro Ser Ala Gly Pro Leu Pro
213          485          490          495
214 Ala Ser Arg Cys Ser Glu Pro Cys Leu Gln Asn Glu Val Lys Ser Val
215          500          505          510
216 Gln Pro Gly Glu Val Cys Cys Trp Leu Cys Ile Pro Cys Gln Pro Tyr
217          515          520          525
218 Glu Tyr Arg Leu Asp Glu Phe Thr Cys Ala Asp Cys Gly Leu Gly Tyr
219          530          535          540
220 Trp Pro Asn Ala Ser Leu Thr Gly Cys Phe Glu Leu Pro Gln Glu Tyr
221 545          550          555          560
222 Ile Arg Trp Gly Asp Ala Trp Ala Val Gly Pro Val Thr Ile Ala Cys
223          565          570          575
224 Leu Gly Ala Leu Ala Thr Leu Phe Val Leu Gly Val Phe Val Arg His
225          580          585          590
226 Asn Ala Thr Pro Val Val Lys Ala Ser Gly Arg Glu Leu Cys Tyr Ile
227          595          600          605
228 Leu Leu Gly Gly Val Phe Leu Cys Tyr Cys Met Thr Phe Ile Phe Ile
229          610          615          620
230 Ala Lys Pro Ser Thr Ala Val Cys Thr Leu Arg Arg Leu Gly Leu Gly
231 625          630          635          640
232 Thr Ala Phe Ser Val Cys Tyr Ser Ala Leu Leu Thr Lys Thr Asn Arg
233          645          650          655
234 Ile Ala Arg Ile Phe Gly Gly Ala Arg Glu Gly Ala Gln Arg Pro Arg
235          660          665          670
236 Phe Ile Ser Pro Ala Ser Gln Val Ala Ile Cys Leu Ala Leu Ile Ser
237          675          680          685
238 Val Gln Leu Leu Ile Val Val Ala Trp Leu Val Val Glu Ala Pro Gly
239          690          695          700
240 Thr Gly Lys Glu Thr Ala Pro Glu Arg Arg Glu Val Val Thr Leu Arg
241 705          710          715          720
242 Cys Asn His Arg Asp Ala Ser Met Leu Gly Ser Leu Ala Tyr Asp Val
243          725          730          735
244 Leu Leu Ile Ala Leu Cys Thr Leu Tyr Ala Phe Lys Thr Arg Lys Cys
245          740          745          750
246 Pro Glu Asn Phe Asn Glu Ala Lys Phe Ile Gly Phe Thr Met Tyr Thr
247          755          760          765
248 Thr Cys Ile Ile Trp Leu Ala Phe Leu Pro Ile Phe Tyr Val Thr Ser
249          770          775          780
250 Ser Asp Tyr Arg Val Gln Thr Thr Thr Met Cys Val Ser Val Ser Leu
251 785          790          795          800
252 Ser Gly Ser Val Val Leu Gly Cys Leu Phe Ala Pro Lys Leu His Ile

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VERIFICATION SUMMARY

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PATENT APPLICATION: **US/10/526,978**

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Input Set : **A:\Ms0023p.txt**

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L:18 M:270 C: Current Application Number differs, Replaced Current Application No

L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date